2017-00187 - Differential model reduction by \((\min,+)\) tropical algebra constraint solving

Level of qualifications required : Graduate degree or equivalent
Fonction : PhD Position

About the research centre or Inria department

Located at the heart of the main national research and higher education cluster, member of the Université Paris Saclay, a major actor in the French Investments for the Future Programme (Idex, LabEx, IRT, Equipex) and partner of the main establishments present on the plateau, the centre is particularly active in three major areas: data and knowledge; safety, security and reliability; modelling, simulation and optimisation (with priority given to energy).

The 450 researchers and engineers from Inria and its partners who work in the research centre’s 31 teams, the 100 research support staff members, the high-level equipment at their disposal (image walls, high-performance computing clusters, sensor networks), and the privileged relationships with prestigious industrial partners, all make Inria Saclay Île-de-France a key research centre in the local landscape and one that is oriented towards Europe and the world.

Context

This thesis subject is offered at Inria Saclay IdF (https://www.inria.fr/en/centre/saclay) in the LIFEWARE project-team (http://lifeware.inria.fr), in the framework of the ANR-DFG SYMBIONT project on "Symbolic Methods for Biological Networks" (https://www.symbiont-project.org/).

The LIFEWARE team works in computational systems biology and develops the Biochemical Abstract Machine (BIOCHAM http://lifeware.inria.fr/biocham) software for modeling, analyzing and now synthesizing biochemical reaction networks (CRNs) using methods from fundamental computer science and mathematics. The software developments are expected to be integrated in BIOCHAM.

The thesis will be supervised by François Fages who has supervised 29 Ph.D. theses in his career, the last defended in May 2016, and who currently is 100% available with no Ph.D. student.

Assignment

Model reduction is a central topic in differential dynamical systems theory, for reducing the complexity of differential equations, finding important parameters, and developing multi-scale models for instance. While perturbation theory is a standard mathematical tool to analyze the different time scales of a dynamical system, and decompose the system according to the fast-slow dynamics, in the domain of computational systems biology, the biochemical reaction networks under consideration involve several tenths of variables which appeal for automated reasoning methods.

Tropicalization is a mathematical method for analysing polynomial equations in the \((\min,+)\) or \((\max,+)\) semiring. The intuitive idea is to reason on the orders of magnitude of concentrations and kinetic parameters which are expressed by the logarithm of their values, hence the \((\min,+)\) algebra for polynomials. We have shown that Constraint Logic Programming can be used to efficiently solve tropical equation problems for identifying the dominating monomials and reactions in a network, and then simplifying the dynamics in the different regions of the phase space corresponding to different regimes (i.e. Quasi Steady State or Quasi Equilibrium assumptions).

The goal of this Thesis, in collaboration with our partners of the ANR-DFG project SYMBIONT, is to generalize our tropical equation solver based on constraint programming to new stability constraints, and to develop the model reduction method by searching for precise approximation results (e.g. using Tikhonov theorem) at least in simple examples like the Michaelis-Menten enzymatic reaction. The method will be implemented in our modeling environment Biocham, evaluated on mcell systems biology models from BioModels repository, and compared to other symbolic methods developed in SYMBIONT.

Main activities

Our first results on this subject have been published in:


The planned work will consist in:

- Generalizing the CLP solver to handle inequalities instead of equalities, with accurate translation from polynomials to \((\min,+)\) intervals;
- Developing correctness criteria for the simplified system (e.g. stability conditions of Tikhonov theorem);
- Implementing the model reduction method and evaluating it on models from the BioModels repository.

Further reading:


General Information

- Theme/Domain : Algorithmics, Computer Algebra and Cryptology
- Scientific computing (BAP E)
- Inria Center : CNRS Saclay - Île-de-France
- Starting date : 2018-09-03
- Duration of contract : 3 years
- Deadline to apply : 2018-08-31

Contacts

- Inria Team : LIFEWARE
- Recruiter : Fages Francois / francois.fages@inria.fr

About Inria

Inria, the French National Institute for computer science and applied mathematics, promotes "scientific excellence for technology transfer and society". Graduates from the world’s top universities, Inria’s 2,700 employees rise to the challenges of digital sciences. With its open, agile model, Inria is able to explore original approaches with its partners in industry and academia and provide an efficient response to the multidisciplinary and application challenges of the digital transformation. Inria is the source of many innovations that add value and create jobs.

The keys to success

We are seeking a highly talented and motivated candidate not afraid by crossing disciplinary frontiers.

Conditions for application

Defence Security :

This position is likely to be situated in a restricted area (ZRR), as defined in Decree No. 2011-1425 relating to the protection of national scientific and technical potential (PPST). Authorisation to enter an area is granted by the director of the unit, following a favourable Ministerial decision, as defined in the decree of 3 July 2012 relating to the PPST. An unfavourable Ministerial decision in respect of a position situated in a ZRR would result in the cancellation of the appointment.

Recruitment Policy :

As part of its diversity policy, all Inria positions are accessible to people with disabilities.

Warning :

You must enter your e-mail address in order to save your application to Inria. Applications must be submitted online on the Inria website. Processing of applications sent from other channels is not guaranteed.
Skills

This subject requires common and basic knowledge on differential equations, linear algebra, algorithms and programming.

There is no specific prerequisite for this internship. However, some specific knowledge on either the algebra (max,+), constraint programming, control theory, or computational systems biology will be a plus.

Benefits package

- Subsidised catering service
- Partially-reimbursed public transport
- Social security
- Paid leave
- Flexible working hours
- Sports facilities

Remuneration

Monthly gross salary: 1.982 euros (1st and 2nd year) - 2.085 euros (3rd year)